

FIGURE 1

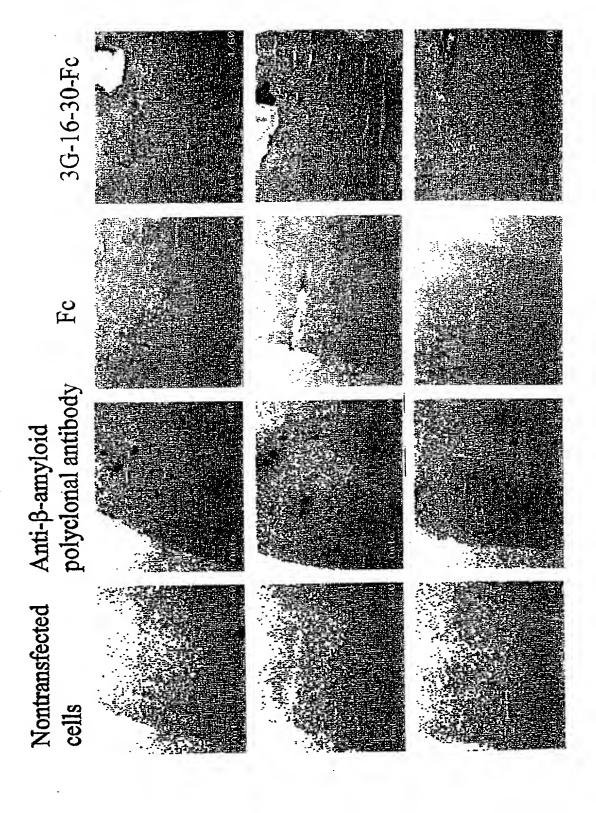


FIGURE 2

#### BssHII-Spe-BamHI converter:

DI215 BSSHII SPEI BAMHI
CGCGCTTCAGAAGAACTAGTG
GAAGTCTTCTTGATCACCTAG DI216
A R F R R T S A S

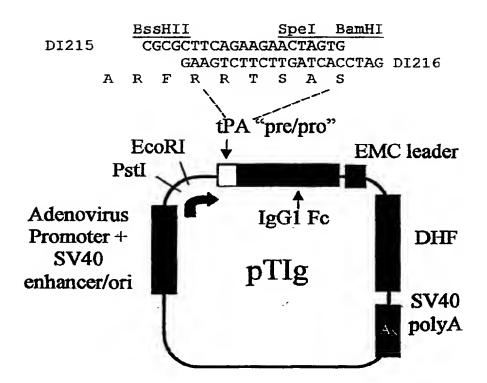


FIGURE 4

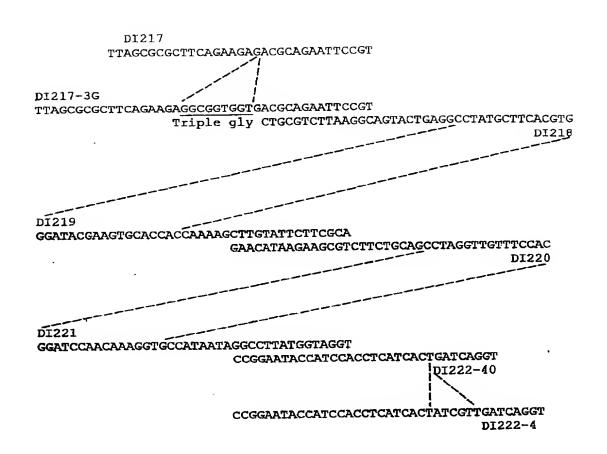


FIGURE 5

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 D A E F R H D S G Y E V H H Q K L V F F A GACGCAGAATTCCGTCATGACTCCGGATACGAAGTGCACCACAAAAGCTTGTATTCTTCGCA EcoRI BspHI BspEI ApalI HinDIII

22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 E D V G S N K G A I I G L M V G G V V I A GAAGACGTCGGATCCAACAAAGGTGCCATAATAGGCCTTATGGTAGGTGGAGTAGTGATAGCA AatII Bamhi Stul

# $\beta$ -amyloid fragments made as IgG1 F<sub>c</sub> fusions:

- LVFFA
- LVFFL
- 16-30 All cloned N-terminal
  10-25 or following triple gly
- 1-29
- 1-40
- 1-42
- also control construct of IgG1 F<sub>c</sub>
   (no β-amyloid sequence)
- Overlapping complimentary oligonucleotides for pentapeptides:

GGGLVFFA:

5′

DI223

CGCGCTTCAGAAGAGGCGGTGGTCTTGTATTCTTCGCAA

GAAGTCTTCTCCGCCACCAGAACATAAGAAGCGTTGATC D1224

BssHII SpeI 5'

LVFFA:

**DI225** 

CGCGCTTCAGAAGACTTGTATTCTTCGCAA

GAAGTCTTCTGAACATAAGAAGCGTTGATC DI226

GGGLVFFL:

DI227

CGCGCTTCAGAAGAGGCGGTGGTCTTGTATTCTTCCTTA
GAAGTCTTCTCCGCCACCAGAACATAAGAAGGAATGATC D1228

FIGURE 7A

LVFFL:

DI229 CGCGCTTCAGAAGACTTGTATTCTTCCTTA
GAAGTCTTCTGAACATAAGAAGGAATGATC DI230

• PCR primers for longer fragments

 $\beta$ -amyloid 1-29 oligos

For 5' use oligos 217 and 217-3G

DI-231 TGGACTAGTACCTTTGTTGGATCCGAC

 $\beta$ -amyloid 10-25 oligos

DI-232 TTAGCGCGCTTCAGAAGATACGAAGTGCACCACCAA

DI-232-3G
TTAGCGCGCTTCAGAAGAGGCGGTGGTTACGAAGTGCACCACCAA

DI-233 TGGACTAGTTCCGACGTCTTCTGCGAA

 $\beta$ -amyloid 16-30 oligos

DI-234 TTAGCGCGCTTCAGAAGAAAGCTTGTATTCTTCGCA

DI-234-3G
TTAGCGCGCTTCAGAAGAGGCGGTGGTAAGCTTGTATTCTTCGCA

DI-235 TGGACTAGTGGCACCTTTGTTGGATCC

### FIGURE 7B

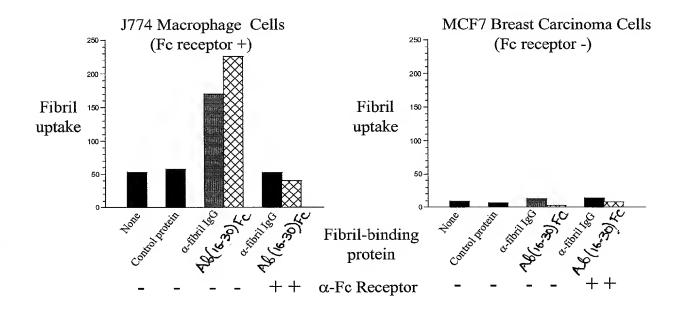


FIGURE 8

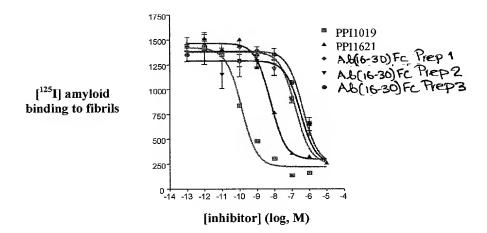
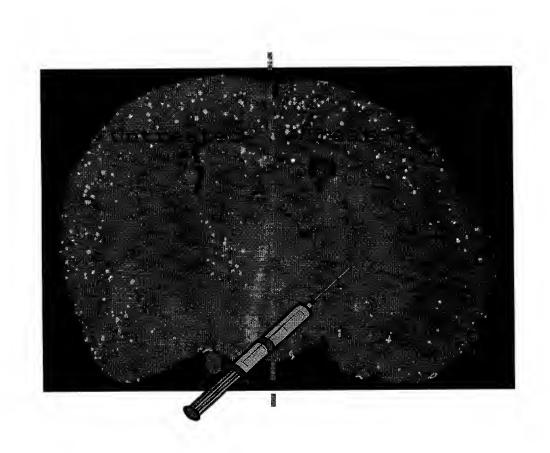


FIGURE 9



ATGGATGCAATGAAGAGAGGGCTCTGCTGTGTGCTGCTGCTGTGTGGAGC AGTCTTCGTTAAGCTTGTATTCTTCGCAGAAGACGTCGGATCGAACAAAG GTGCCGAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCA GCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTCTTCCCCCCAAAACCC AAGGACACCCTCATGATATCCCGGACCCCTGAGGTCACATGCGTGGTGGT GGACGTGAGCCACGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACG GCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAGTACAA CAGCACGTACCGGGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGC TGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAACAAAGCCCTCCCAGCC CCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCAC AGGTGTACACCCTGCCCCCATCCCGGGATGAGCTGACCAAGAACCAGGTC AGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAGCGACATCGCCGTGGA GTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCC GTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGAC AAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTCCGTGATGCATGA GGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTA **AATGA** 

